

=> dis his

(FILE 'HOME' ENTERED AT 11:33:49 ON 12 JAN 2006)

FILE 'MEDLINE, BIOSIS, CAPLUS, EMBASE, SCISEARCH' ENTERED AT 11:34:07 ON 12 JAN 2006

L1 668 S ((PRIMER EXTENS? OR MINISEQUENCING) (20A) (ARRAY? OR BLOT? OR
L2 272 DUP REM L1 (396 DUPLICATES REMOVED)
L3 123 S L2 AND PY<2002
L4 11 S L3 AND (DDGTP OR DDCTP OR DDATP OR DDTTP OR DIDEOXY?)

FILE 'STNGUIDE' ENTERED AT 11:40:16 ON 12 JAN 2006

L5 0 S ((PRIMER EXTENS? OR MINISEQUENCING OR SNUPE OR SNAPSHOT OR AP

FILE 'MEDLINE, BIOSIS, CAPLUS, EMBASE, SCISEARCH' ENTERED AT 11:51:08 ON 12 JAN 2006

L6 751 S ((PRIMER EXTENS? OR MINISEQUENCING OR SNUPE OR SNAPSHOT OR A
L7 311 DUP REM L6 (440 DUPLICATES REMOVED)
L8 149 S L7 AND PY<2002
L9 27 S L8 AND (DDNTP? OR DDGTP OR DDCTP OF DDATP OF DDTTP OR DIDEOX

FILE 'STNGUIDE' ENTERED AT 12:00:20 ON 12 JAN 2006

FILE 'MEDLINE, BIOSIS, CAPLUS, EMBASE, SCISEARCH' ENTERED AT 12:19:47 ON 12 JAN 2006

FILE 'MEDLINE, BIOSIS, CAPLUS, EMBASE, SCISEARCH' ENTERED AT 12:22:55 ON 12 JAN 2006

L10 751 S ((PRIMER EXTENS? OR MINISEQUENCING OR SNUPE OR SNAPSHOT OR A
L11 311 DUP REM L10 (440 DUPLICATES REMOVED)
L12 149 S L11 AND PY<2002

FILE 'HOME' ENTERED AT 12:56:28 ON 12 JAN 2006

=> FIL MEDLINE BIOSIS CAPLUS EMBASE SCISEARCH
COST IN U.S. DOLLARS

SINCE FILE	TOTAL
ENTRY	SESSION
0.21	0.21

FULL ESTIMATED COST

FILE 'MEDLINE' ENTERED AT 12:56:40 ON 12 JAN 2006

FILE 'BIOSIS' ENTERED AT 12:56:40 ON 12 JAN 2006

Copyright (c) 2006 The Thomson Corporation

FILE 'CAPLUS' ENTERED AT 12:56:40 ON 12 JAN 2006

USE IS SUBJECT TO THE TERMS OF YOUR STN CUSTOMER AGREEMENT.

PLEASE SEE "HELP USAGETERMS" FOR DETAILS.

COPYRIGHT (C) 2006 AMERICAN CHEMICAL SOCIETY (ACS)

FILE 'EMBASE' ENTERED AT 12:56:40 ON 12 JAN 2006

Copyright (c) 2006 Elsevier B.V. All rights reserved.

FILE 'SCISEARCH' ENTERED AT 12:56:40 ON 12 JAN 2006

Copyright (c) 2006 The Thomson Corporation

=> (primer estens? or minisequencing or apex or apse or snupe or snapshot) and (a;
(PRIMER IS NOT A RECOGNIZED COMMAND

The previous command name entered was not recognized by the system.

For a list of commands available to you in the current file, enter

"HELP COMMANDS" at an arrow prompt (=>).

=> s (primer estens? or minisequencing or apex or apse or snupe or snapshot) and
L1 474 (PRIMER ESTENS? OR MINISEQUENCING OR APEX OR APSE OR SNUPE OR
SNAPSHOT) AND (ARRAY? OR BLOT? OR MEMBRANE? OR CHIP?) AND (SNP?
OR MUTATION? OR SUBSTITUTION? OR GENOTYPE? OR ALLELE? OR HAPLOTY
PE?)

=> s l1 and (ddntp? or ddatp or ddttp or dideoxy? or (chain (w) terminat?))

L2 55 L1 AND (DDNTP? OR DDATP OR DDTTP OR DIDEOXY? OR (CHAIN (W) TERMI
NAT?))

=> dup rem 12

PROCESSING COMPLETED FOR L2

L3 30 DUP REM L2 (25 DUPLICATES REMOVED)

=> s l3 and py<2003

L4 20 L3 AND PY<2003

=> s (primer extends? or minisequencing or apex or apse or snupe or snapshot) and
L5 1843 (PRIMER EXTENS? OR MINISEQUENCING OR APEX OR APSE OR SNUPE OR
SNAPSHOT) AND (ARRAY? OR BLOT? OR MEMBRANE? OR CHIP?) AND (SNP?
OR MUTATION? OR SUBSTITUTION? OR GENOTYPE? OR ALLELE? OR HAPLOTY
PE?)

=> s l5 and (ddntp? or ddatp or ddttp or dideoxy? or (chain (w) terminat?))

L6 110 L5 AND (DDNTP? OR DDATP OR DDTTP OR DIDEOXY? OR (CHAIN (W)
TERMINAT?))

=> dup rem l6

PROCESSING COMPLETED FOR L6

L7 65 DUP REM L6 (45 DUPLICATES REMOVED)

=> dis l7 1-65 bib, ab

=>